

10/566054

-1-

AP20 Rec'd OCT 25 JAN 2006
SEQUENCE LISTING

<110> Ambrose, Helen Jean
March, Ruth

<120> USE OF POLYMORPHISMS IN HUMAN OATP-C ASSOCIATED WITH AN EFFECT ON STATIN
PHARMACOKINETICS IN HUMANS IN STATIN THERAPY

<130> 06275-492US1

<150> PCT/GB2004/003236

<151> 2004-07-26

<150> GB 0317592.4

<151> 2003-07-26

<160> 5

<170> PatentIn version 3.1

<210> 1

<211> 691

<212> PRT

<213> Homo Sapiens

<400> 1

Met Asp Gln Asn Gln His Leu Asn Lys Thr Ala Glu Ala Gln Pro Ser
1 5 10 15

Glu Asn Lys Lys Thr Arg Tyr Cys Asn Gly Leu Lys Met Phe Leu Ala
20 25 30

Ala Leu Ser Leu Ser Phe Ile Ala Lys Thr Leu Gly Ala Ile Ile Met
35 40 45

Lys Ser Ser Ile Ile His Ile Glu Arg Arg Phe Glu Ile Ser Ser Ser
50 55 60

Leu Val Gly Phe Ile Asp Gly Ser Phe Glu Ile Gly Asn Leu Leu Val
65 70 75 80

Ile Val Phe Val Ser Tyr Phe Gly Ser Lys Leu His Arg Pro Lys Leu
85 90 95

Ile Gly Ile Gly Cys Phe Ile Met Gly Ile Gly Gly Val Leu Thr Ala
100 105 110

Leu Pro His Phe Phe Met Gly Tyr Tyr Arg Tyr Ser Lys Glu Thr Asn
115 120 125

Ile Asn Ser Ser Glu Asn Ser Thr Ser Thr Leu Ser Thr Cys Leu Ile
130 135 140

Asn Gln Ile Leu Ser Leu Asn Arg Ala Ser Pro Glu Ile Val Gly Lys
145 150 155 160

Gly Cys Leu Lys Glu Ser Gly Ser Tyr Met Trp Ile Tyr Val Phe Met
165 170 175

Gly Asn Met Leu Arg Gly Ile Gly Glu Thr Pro Ile Val Pro Leu Gly
180 185 190

Leu Ser Tyr Ile Asp Asp Phe Ala Lys Glu Gly His Ser Ser Leu Tyr
195 200 205

Leu Gly Ile Leu Asn Ala Ile Ala Met Ile Gly Pro Ile Ile Gly Phe
210 215 220

Thr Leu Gly Ser Leu Phe Ser Lys Met Tyr Val Asp Ile Gly Tyr Val
225 230 235 240

Asp Leu Ser Thr Ile Arg Ile Thr Pro Thr Asp Ser Arg Trp Val Gly
245 250 255

Ala Trp Trp Leu Asn Phe Leu Val Ser Gly Leu Phe Ser Ile Ile Ser
260 265 270

Ser Ile Pro Phe Phe Phe Leu Pro Gln Thr Pro Asn Lys Pro Gln Lys
275 280 285

Glu Arg Lys Ala Ser Leu Ser Leu His Val Leu Glu Thr Asn Asp Glu
290 295 300

Lys Asp Gln Thr Ala Asn Leu Thr Asn Gln Gly Lys Asn Ile Thr Lys
305 310 315 320

Asn Val Thr Gly Phe Phe Gln Ser Phe Lys Ser Ile Leu Thr Asn Pro
325 330 335

Leu Tyr Val Met Phe Val Leu Leu Thr Leu Leu Gln Val Ser Ser Tyr
340 345 350

Ile Gly Ala Phe Thr Tyr Val Phe Lys Tyr Val Glu Gln Gln Tyr Gly
355 360 365

Gln Pro Ser Ser Lys Ala Asn Ile Leu Leu Gly Val Ile Thr Ile Pro
370 375 380

Ile Phe Ala Ser Gly Met Phe Leu Gly Gly Tyr Ile Ile Lys Lys Phe
385 390 395 400

Lys Leu Asn Thr Val Gly Ile Ala Lys Phe Ser Cys Phe Thr Ala Val
405 410 415

Met Ser Leu Ser Phe Tyr Leu Leu Tyr Phe Phe Ile Leu Cys Glu Asn
420 425 430

Lys Ser Val Ala Gly Leu Thr Met Thr Tyr Asp Gly Asn Asn Pro Val
435 440 445

Thr Ser His Arg Asp Val Pro Leu Ser Tyr Cys Asn Ser Asp Cys Asn
450 455 460

Cys Asp Glu Ser Gln Trp Glu Pro Val Cys Gly Asn Asn Gly Ile Thr
465 470 475 480

Tyr Ile Ser Pro Cys Leu Ala Gly Cys Lys Ser Ser Ser Gly Asn Lys
485 490 495

Lys Pro Ile Val Phe Tyr Asn Cys Ser Cys Leu Glu Val Thr Gly Leu
500 505 510

Gln Asn Arg Asn Tyr Ser Ala His Leu Gly Glu Cys Pro Arg Asp Asp
515 520 525

Ala Cys Thr Arg Lys Phe Tyr Phe Phe Val Ala Ile Gln Val Leu Asn
530 535 540

Leu Phe Phe Ser Ala Leu Gly Gly Thr Ser His Val Met Leu Ile Val
545 550 555 560

Lys Ile Val Gln Pro Glu Leu Lys Ser Leu Ala Leu Gly Phe His Ser
565 570 575

Met Val Ile Arg Ala Leu Gly Gly Ile Leu Ala Pro Ile Tyr Phe Gly
580 585 590

Ala Leu Ile Asp Thr Thr Cys Ile Lys Trp Ser Thr Asn Asn Cys Gly
595 600 605

Thr Arg Gly Ser Cys Arg Thr Tyr Asn Ser Thr Ser Phe Ser Arg Val
610 615 620

Tyr Leu Gly Leu Ser Ser Met Leu Arg Val Ser Ser Leu Val Leu Tyr
625 630 635 640

Ile Ile Leu Ile Tyr Ala Met Lys Lys Lys Tyr Gln Glu Lys Asp Ile
645 650 655

Asn Ala Ser Glu Asn Gly Ser Val Met Asp Glu Ala Asn Leu Glu Ser
660 665 670

Leu Asn Lys Asn Lys His Phe Val Pro Ser Ala Gly Ala Asp Ser Glu
675 680 685

Thr His Cys

690

<210> 2
<211> 2050
<212> DNA
<213> Homo Sapiens

<400> 2
atctcagaga ttttatttgt attcatttaa tataaattaa ctgctctaaa atttataata 60
tgcaaatatc atacaattaa tctaattagg tgttgaatct ataatgtgcc aggcattatg 120
taaggcactt tacatacact aaatctttat tccaaatata gacttcttac tttatagatg 180
agtgcactga tgctcagaaa tggtaaataa cctactgatg tttatactgc tggcaggtag 240
cagagacata tcggcattta agtctttcag acttcaaagg ccatgatatt tcatcagagc 300
tgtgatagcc gttcctgaaa aaaatatcag ctgattcttt aaatcaattt ttgtcatcta 360
actgatgcgt ggctgttagc ataatttga tcttgaaaga tgttttgcaa catctttccc 420
ctggtgtact cttgtttttc catgatccca caaatgagc agtctaatta tttacacaat 480
taggaagaga aaaggggcac agagaatgct ctttgacctc tgaaaatatt ggagaatttt 540
acaactggca ccttttagctc aggattataa aggttgtag ttagtttgta ctgttttatc 600
ttcattgtat ataatatata tattagtctc caaacatgtt gatgtgtttt caatgaaatg 660
gatgtctgag gagaaaacca ttagcctgag aaaacccaaa ctgtattccc atttgtgaata 720
aaaggaagtc cataaaaatg atggaaaatg ttctgcattc ctgttatgat atcaaaaatc 780
ggcagtacat gaaaattttt caaagtgtt atttaacagg cataatcttt ggtctcctga 840
gccagaatct gctgggtatg ggactggatt gctattttga caactcgcca gtagattctt 900
actcagcaga gtatttgga gcttactct aatattttgg ccttggttct acatttctca 960
gttctgcaca gtcattcttc cctctacac tactctttag tttgtctcat gattccaata 1020
ctctcaataa ttaaccaaga atagaactaa tcaatcagat aactgtggca cagacatcaa 1080
atacattttg ctgcaaccat atcaacaaat gtcccatgaa tgataagggg taaccatatt 1140
ctcatatatg catctcaca ttaccacata tatatatgtg catatgtgta tacaggtaaa 1200
agtgtgtata tatgtatata tgtatgtttg tgtgtatata catacatata tcttcacact 1260
tttctgaaat atatatattt atgtgagaga agggctctgta ctttatttca gaagagagct 1320
taatgtccaa ggtataattg agagtctaaa atgtttgagt tattgaatta attaaacttc 1380
atctctactc aagaaaactt ttaactgagt taagctcttc ctttctccac aagtcaagtc 1440
aataaaagga aactgtgata ttaataattc tttcctgttt tgatgtaaag aatctatcgc 1500
ataaagcagt ctttaattttc atcattcaga aaaatggtct tgcagttaat tgggactctc 1560
ttattccagg tggatatctc agtctccata cataccacgt tagaaccata cttatgtacc 1620
aagcaaagag ggtatatattt aattttttaa tgccaatgta acctgtaggc atatttttta 1680
tttgtcttaa attatttcct atttggaggt tttaaatacc tggataaatt tattgtactc 1740

atatttttaa agaaaaaat cttatgccac caacttaatt gaataaacia gtaaaagcca 1800
ttcccaaaag taaggtttac ttgttaagat taacaaaaaa taatgtgaga attctgagaa 1860
atataatctt taaatattgg caactggagt gaactcttaa aactaactag gttttatatg 1920
tttgactaga gcaatgacat aataaggtgg ttaatcatca ctggacttgt tttcaaaaag 1980
ccaactactt taagaggaat aaaggggtga cttgttgagc ttgctgtagg attctaaatc 2040
caggttaagaa 2050

<210> 3
<211> 2830
<212> DNA
<213> Homo Sapiens

<400> 3
cggacgcgtg ggcggacgcg tgggtcgccc acgcgtccga cttgttgagc ttgctgtagg 60
attctaaatc caggtgattg tttcaaactg agcatcaaca aaaaaaacat ttgtatgata 120
tctatatttc aatcatggac caaatcaac atttgaataa aacagcagag gcacaacctt 180
cagagaataa gaaaacaaga tactgcaatg gattgaagat gttcttgagc gctctgtcac 240
tcagctttat tgctaagaca ctaggtgcaa ttattatgaa aagttccatc attcatatag 300
aacggagatt tgagatatcc tcttctcttg ttggttttat tgacggaagc tttgaaattg 360
gaaatttgct tgtgattgta tttgtgagtt actttggatc caaactacat agaccaaagt 420
taattggaat cggttgtttc attatgggaa ttggaggtgt tttgactgct ttgccacatt 480
tcttcatggg atattacagg tattctaaag aaactaatat cgattcatca gaaaattcaa 540
catcgacctt atccacttgt ttaattaatc aaattttatc actcaataga gcatcacctg 600
agatagtggg aaaagggtgt ttaaaggaat ctgggtcata catgtggata tatgtgttca 660
tgggtaatat gcttcgtgga ataggggaga ctcccatagt accattgggg ctttcttaca 720
ttgatgattt cgctaaagaa ggacattctt ctttgtattt aggtatattg aatgcaatag 780
caatgattgg tccaatcatt ggctttaccg tgggatctct gttttctaaa atgtacgtgg 840
atattggata ttagatcta agcactatca ggataactcc tactgattct cgatggggtg 900
gagcttggtg gcttaatttc cttgtgtctg gactattctc cattatttct tccataccat 960
tctttttctt gcccctaaact ccaaataaac cacaaaaaga aagaaaagct tcaactgtctt 1020
tgcatgtgct ggaaacaaat gatgaaaagg atcaaacagc taatttgacc aatcaaggaa 1080
aaaatattac caaaaatgtg actgggtttt tccagtcttt taaaagcatc cttactaatc 1140
ccctgtatgt tatgtttgtg cttttgacgt tgttacaagt aagcagctat attgggtgctt 1200
ttacttatgt cttcaaatac gtagagcaac agtatggtca gccttcatct aaggctaaca 1260
tcttattggg agtcataacc atacctatct ttgcaagtgg aatgttttta ggaggatata 1320
tcattaaaaa attcaaactg aacaccgttg gaattgccaa attctcatgt tttactgctg 1380
tgatgtcatt gtccttttac ctattatatt ttttcatact ctgtgaaaac aaatcagttg 1440

ccggactaac catgacctat gatggaaata atccagtgc atctcataga gatgtaccac 1500
tttcttattg caactcagac tgcaattgtg atgaaagtca atgggaacca gtctgtggaa 1560
acaatggaat aacttacatc tcacctgtc tagcagggtg caaatcttca agtggcaata 1620
aaaagcctat agtgttttac aactgcagtt gtttggaagt aactgggtctc cagaacagaa 1680
attactcagc ccatttggtt gaatgcccaa gagatgatgc ttgtacaagg aaattttact 1740
tttttggtgc aatacaagtc ttgaatttat ttttctctgc acttgagggc acctcacatg 1800
tcatgctgat tgttaaaatt gttcaacctg aattgaaatc acttgactg ggtttccact 1860
caatgggtat acgagcacta ggaggaattc tagctccaat atattttggg gctctgattg 1920
atacaacgtg tataaagtgg tccaccaaca actgtggcac acgtgggtca tgtaggacat 1980
ataattccac atcattttca aggtgtctact tgggcttgtc ttcaatgta agagtctcat 2040
cacttgtttt atatattata ttaatttatg ccatgaagaa aaaatatcaa gagaaagata 2100
tcaatgcac agaaaatgga agtgtcatgg atgaagcaaa cttagaatcc ttaaataaaa 2160
ataaacattt tgtcccttct gctggggcag atagtgaac acattgttaa ggggagaaaa 2220
aaagccactt ctgcttctgt gtttccaaac agcattgcat tgattcagta agatgttatt 2280
tttgaggagt tcctggtcct ttcactaaga atttccacat cttttatggt ggaagtataa 2340
ataagcctat gaacttataa taaaacaaac tgtaggtaga aaaaatgaga gtactcattg 2400
ttacattata gctacatatt tgtggttaag gttagactat atgatccata caaattaaag 2460
tgagagacat gggtactgtg taataaaaga aaaaatactt gttcaggtaa ttctaattct 2520
taataaaaca aatgagtatc atacaggtag aggttaaaaa ggaggagcta gattcatatc 2580
ctaagtaaag agaatgcct agtgtctatt ttattaaaca aacaaacaca gagtttgaac 2640
tataatacta aggcctgaag tctagcttgg atatatgcta caataatata tgttactcac 2700
ataaaattat atatttcaca gactttatca atgtataatt aacaattatc ttgtttaagt 2760
aaatttagaa tacatttaag tattgtggaa gaaataaaga cattccaata ttgcaaaaaa 2820
aaaaaaaaa 2830

<210> 4
<211> 691
<212> PRT
<213> Homo Sapiens

<400> 4

Met Asp Gln Asn Gln His Leu Asn Lys Thr Ala Glu Ala Gln Pro Ser
1 5 10 15

Glu Asn Lys Lys Thr Arg Tyr Cys Asn Gly Leu Lys Met Phe Leu Ala
20 25 30

Ala Leu Ser Leu Ser Phe Ile Ala Lys Thr Leu Gly Ala Ile Ile Met
35 40 45

Lys Ser Ser Ile Ile His Ile Glu Arg Arg Phe Glu Ile Ser Ser Ser
 50 55 60
 Leu Val Gly Phe Ile Asp Gly Ser Phe Glu Ile Gly Asn Leu Leu Val
 65 70 75 80
 Ile Val Phe Val Ser Tyr Phe Gly Ser Lys Leu His Arg Pro Lys Leu
 85 90 95
 Ile Gly Ile Gly Cys Phe Ile Met Gly Ile Gly Gly Val Leu Thr Ala
 100 105 110
 Leu Pro His Phe Phe Met Gly Tyr Tyr Arg Tyr Ser Lys Glu Thr Asn
 115 120 125
 Ile Asp Ser Ser Glu Asn Ser Thr Ser Thr Leu Ser Thr Cys Leu Ile
 130 135 140
 Asn Gln Ile Leu Ser Leu Asn Arg Ala Ser Pro Glu Ile Val Gly Lys
 145 150 155 160
 Gly Cys Leu Lys Glu Ser Gly Ser Tyr Met Trp Ile Tyr Ala Phe Met
 165 170 175
 Gly Asn Met Leu Arg Gly Ile Gly Glu Thr Pro Ile Val Pro Leu Gly
 180 185 190
 Leu Ser Tyr Ile Asp Asp Phe Ala Lys Glu Gly His Ser Ser Leu Tyr
 195 200 205
 Leu Gly Ile Leu Asn Ala Ile Ala Met Ile Gly Pro Ile Ile Gly Phe
 210 215 220
 Thr Leu Gly Ser Leu Phe Ser Lys Met Tyr Val Asp Ile Gly Tyr Val
 225 230 235 240
 Asp Leu Ser Thr Ile Arg Ile Thr Pro Thr Asp Ser Arg Trp Val Gly
 245 250 255
 Ala Trp Trp Leu Asn Phe Leu Val Ser Gly Leu Phe Ser Ile Ile Ser
 260 265 270
 Ser Ile Pro Phe Phe Phe Leu Pro Gln Thr Pro Asn Lys Pro Gln Lys
 275 280 285
 Glu Arg Lys Ala Ser Leu Ser Leu His Val Leu Glu Thr Asn Asp Glu
 290 295 300
 Lys Asp Gln Thr Ala Asn Leu Thr Asn Gln Gly Lys Asn Ile Thr Lys
 305 310 315 320

Asn Val Thr Gly Phe Phe Gln Ser Phe Lys Ser Ile Leu Thr Asn Pro
325 330 335

Leu Tyr Val Met Phe Val Leu Leu Thr Leu Leu Gln Val Ser Ser Tyr
340 345 350

Ile Gly Ala Phe Thr Tyr Val Phe Lys Tyr Val Glu Gln Gln Tyr Gly
355 360 365

Gln Pro Ser Ser Lys Ala Asn Ile Leu Leu Gly Val Ile Thr Ile Pro
370 375 380

Ile Phe Ala Ser Gly Met Phe Leu Gly Gly Tyr Ile Ile Lys Lys Phe
385 390 395 400

Lys Leu Asn Thr Val Gly Ile Ala Lys Phe Ser Cys Phe Thr Ala Val
405 410 415

Met Ser Leu Ser Phe Tyr Leu Leu Tyr Phe Phe Ile Leu Cys Glu Asn
420 425 430

Lys Ser Val Ala Gly Leu Thr Met Thr Tyr Asp Gly Asn Asn Pro Val
435 440 445

Thr Ser His Arg Asp Val Pro Leu Ser Tyr Cys Asn Ser Asp Cys Asn
450 455 460

Cys Asp Glu Ser Gln Trp Glu Pro Val Cys Gly Asn Asn Gly Ile Thr
465 470 475 480

Tyr Ile Ser Pro Cys Leu Ala Gly Cys Lys Ser Ser Ser Gly Asn Lys
485 490 495

Lys Pro Ile Val Phe Tyr Asn Cys Ser Cys Leu Glu Val Thr Gly Leu
500 505 510

Gln Asn Arg Asn Tyr Ser Ala His Leu Gly Glu Cys Pro Arg Asp Asp
515 520 525

Ala Cys Thr Arg Lys Phe Tyr Phe Phe Val Ala Ile Gln Val Leu Asn
530 535 540

Leu Phe Phe Ser Ala Leu Gly Gly Thr Ser His Val Met Leu Ile Val
545 550 555 560

Lys Ile Val Gln Pro Glu Leu Lys Ser Leu Ala Leu Gly Phe His Ser
565 570 575

Met Val Ile Arg Ala Leu Gly Gly Ile Leu Ala Pro Ile Tyr Phe Gly

580 585 590

Ala Leu Ile Asp Thr Thr Cys Ile Lys Trp Ser Thr Asn Asn Cys Gly
595 600 605

Thr Arg Gly Ser Cys Arg Thr Tyr Asn Ser Thr Ser Phe Ser Arg Val
610 615 620

Tyr Leu Gly Leu Ser Ser Met Leu Arg Val Ser Ser Leu Val Leu Tyr
625 630 635 640

Ile Ile Leu Ile Tyr Ala Met Lys Lys Lys Tyr Gln Glu Lys Asp Ile
645 650 655

Asn Ala Ser Glu Asn Gly Ser Val Met Asp Glu Ala Asn Leu Glu Ser
660 665 670

Leu Asn Lys Asn Lys His Phe Val Pro Ser Ala Gly Ala Asp Ser Glu
675 680 685

Thr His Cys
690

<210> 5
<211> 691
<212> PRT
<213> Homo Sapiens

<400> 5

Met Asp Gln Asn Gln His Leu Asn Lys Thr Ala Glu Ala Gln Pro Ser
1 5 10 15

Glu Asn Lys Lys Thr Arg Tyr Cys Asn Gly Leu Lys Met Phe Leu Ala
20 25 30

Ala Leu Ser Leu Ser Phe Ile Ala Lys Thr Leu Gly Ala Ile Ile Met
35 40 45

Lys Ser Ser Ile Ile His Ile Glu Arg Arg Phe Glu Ile Ser Ser Ser
50 55 60

Leu Val Gly Phe Ile Asp Gly Ser Phe Glu Ile Gly Asn Leu Leu Val
65 70 75 80

Ile Val Phe Val Ser Tyr Phe Gly Ser Lys Leu His Arg Pro Lys Leu
85 90 95

Ile Gly Ile Gly Cys Phe Ile Met Gly Ile Gly Gly Val Leu Thr Ala
100 105 110

Leu Pro His Phe Phe Met Gly Tyr Tyr Arg Tyr Ser Lys Glu Thr Asn

115	120	125
Ile Asn Ser Ser Glu Asn Ser Thr Ser Thr Leu Ser Thr Cys Leu Ile		
130	135	140
Asn Gln Ile Leu Ser Leu Asn Arg Ala Ser Pro Glu Ile Val Gly Lys		
145	150	155 160
Gly Cys Leu Lys Glu Ser Gly Ser Tyr Met Trp Ile Tyr Ala Phe Met		
	165	170 175
Gly Asn Met Leu Arg Gly Ile Gly Glu Thr Pro Ile Val Pro Leu Gly		
	180	185 190
Leu Ser Tyr Ile Asp Asp Phe Ala Lys Glu Gly His Ser Ser Leu Tyr		
	195	200 205
Leu Gly Ile Leu Asn Ala Ile Ala Met Ile Gly Pro Ile Ile Gly Phe		
	210	215 220
Thr Leu Gly Ser Leu Phe Ser Lys Met Tyr Val Asp Ile Gly Tyr Val		
	225	230 235 240
Asp Leu Ser Thr Ile Arg Ile Thr Pro Thr Asp Ser Arg Trp Val Gly		
	245	250 255
Ala Trp Trp Leu Asn Phe Leu Val Ser Gly Leu Phe Ser Ile Ile Ser		
	260	265 270
Ser Ile Pro Phe Phe Phe Leu Pro Gln Thr Pro Asn Lys Pro Gln Lys		
	275	280 285
Glu Arg Lys Ala Ser Leu Ser Leu His Val Leu Glu Thr Asn Asp Glu		
	290	295 300
Lys Asp Gln Thr Ala Asn Leu Thr Asn Gln Gly Lys Asn Ile Thr Lys		
	305	310 315 320
Asn Val Thr Gly Phe Phe Gln Ser Phe Lys Ser Ile Leu Thr Asn Pro		
	325	330 335
Leu Tyr Val Met Phe Val Leu Leu Thr Leu Leu Gln Val Ser Ser Tyr		
	340	345 350
Ile Gly Ala Phe Thr Tyr Val Phe Lys Tyr Val Glu Gln Gln Tyr Gly		
	355	360 365
Gln Pro Ser Ser Lys Ala Asn Ile Leu Leu Gly Val Ile Thr Ile Pro		
	370	375 380

Ile Phe Ala Ser Gly Met Phe Leu Gly Gly Tyr Ile Ile Lys Lys Phe
385 390 395 400

Lys Leu Asn Thr Val Gly Ile Ala Lys Phe Ser Cys Phe Thr Ala Val
405 410 415

Met Ser Leu Ser Phe Tyr Leu Leu Tyr Phe Phe Ile Leu Cys Glu Asn
420 425 430

Lys Ser Val Ala Gly Leu Thr Met Thr Tyr Asp Gly Asn Asn Pro Val
435 440 445

Thr Ser His Arg Asp Val Pro Leu Ser Tyr Cys Asn Ser Asp Cys Asn
450 455 460

Cys Asp Glu Ser Gln Trp Glu Pro Val Cys Gly Asn Asn Gly Ile Thr
465 470 475 480

Tyr Ile Ser Pro Cys Leu Ala Gly Cys Lys Ser Ser Ser Gly Asn Lys
485 490 495

Lys Pro Ile Val Phe Tyr Asn Cys Ser Cys Leu Glu Val Thr Gly Leu
500 505 510

Gln Asn Arg Asn Tyr Ser Ala His Leu Gly Glu Cys Pro Arg Asp Asp
515 520 525

Ala Cys Thr Arg Lys Phe Tyr Phe Phe Val Ala Ile Gln Val Leu Asn
530 535 540

Leu Phe Phe Ser Ala Leu Gly Gly Thr Ser His Val Met Leu Ile Val
545 550 555 560

Lys Ile Val Gln Pro Glu Leu Lys Ser Leu Ala Leu Gly Phe His Ser
565 570 575

Met Val Ile Arg Ala Leu Gly Gly Ile Leu Ala Pro Ile Tyr Phe Gly
580 585 590

Ala Leu Ile Asp Thr Thr Cys Ile Lys Trp Ser Thr Asn Asn Cys Gly
595 600 605

Thr Arg Gly Ser Cys Arg Thr Tyr Asn Ser Thr Ser Phe Ser Arg Val
610 615 620

Tyr Leu Gly Leu Ser Ser Met Leu Arg Val Ser Ser Leu Val Leu Tyr
625 630 635 640

Ile Ile Leu Ile Tyr Ala Met Lys Lys Lys Tyr Gln Glu Lys Asp Ile
645 650 655

Asn Ala Ser Glu Asn Gly Ser Val Met Asp Glu Ala Asn Leu Glu Ser
660 665 670

Leu Asn Lys Asn Lys His Phe Val Pro Ser Ala Gly Ala Asp Ser Glu
675 680 685

Thr His Cys
690